



#6

SEQUENCE LISTING

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NILSSON, Magnus  
MORISSON, Veronique

<120> Cysteine Protease Inhibitors

<130> 1718-0195P

<140> US 10/015,186

<141> 2001-11-16

<150> US 60/252,840

<151> 2000-11-17

<150> PCT/GB00/01894

<151> 2000-05-18

<150> GB 9911417.5

<151> 1999-05-18

<160> 4

<170> PatentIn version 3.0

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<212> DNA

<213> Artificial Sequence

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<223> Primer for cDNA of cysteinyl proteinase (Falcipain 2)

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aac aaa tat ctt agt tta aga tct tca aaa cca tta aag aat tct aaa 95  
Asn Lys Tyr Leu Ser Leu Arg Ser Ser Lys Pro Leu Lys Asn Ser Lys  
20 25 30

tat tta tta gat caa atg aat tat gaa gaa gtt ata aaa aaa tat aga 143  
Tyr Leu Leu Asp Gln Met Asn Tyr Glu Glu Val Ile Lys Lys Tyr Arg  
35 40 45

gga gaa gaa aat ttc gat cat gca gct tac gac tgg aga tta cac agt 191  
Gly Glu Glu Asn Phe Asp His Ala Ala Tyr Asp Trp Arg Leu His Ser  
50 55 60

ggt gta aca cct gta aag gat caa aaa aat tgt gga tct tgc tgg gcc 239  
Gly Val Thr Pro Val Lys Asp Gln Lys Asn Cys Gly Ser Cys Trp Ala  
65 70 75

ttt agt agt ata ggt tcc gta gaa tca caa tat gct atc aga aaa aat 287  
Phe Ser Ser Ile Gly Ser Val Glu Ser Gln Tyr Ala Ile Arg Lys Asn  
80 85 90 95

aaa tta ata acc tta agt gaa caa gaa tta gta gat tgt tca ttt aaa 335  
Lys Leu Ile Thr Leu Ser Glu Gln Glu Leu Val Asp Cys Ser Phe Lys  
100 105 110

aat tat ggt tgt aat gga ggt ctc att aat aat gcc ttt gag gat atg 383  
Asn Tyr Gly Cys Asn Gly Gly Leu Ile Asn Asn Ala Phe Glu Asp Met  
115 120 125

att gaa ctt gga ggt ata tgt cca gat ggt gat tat cca tat gtg agt 431  
Ile Glu Leu Gly Gly Ile Cys Pro Asp Gly Asp Tyr Pro Tyr Val Ser  
130 135 140

gat gct cca aat tta tgt aac ata gat aga tgt act gaa aaa tat gga 479  
Asp Ala Pro Asn Leu Cys Asn Ile Asp Arg Cys Thr Glu Lys Tyr Gly  
145 150 155

atc aaa aat tat tta tcc gta cca gat aat aaa tta aaa gaa gca ctt 527  
Ile Lys Asn Tyr Leu Ser Val Pro Asp Asn Lys Leu Lys Glu Ala Leu  
160 165 170 175

aga ttc ttg gga cct att agt att agt gta gcc gta tca gat gat ttt 575  
Arg Phe Leu Gly Pro Ile Ser Ile Ser Val Ala Val Ser Asp Asp Phe  
180 185 190

gct ttt tac aaa gaa ggt att ttc gat gga gaa tgt ggt gat gaa tta	623
Ala Phe Tyr Lys Glu Gly Ile Phe Asp Gly Glu Cys Gly Asp Glu Leu	
195 200 205	

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Asn His Ala Val Met Leu Val Gly Phe Gly Met Lys Glu Ile Val Asn	
210 215 220	

cca tta acc aag aaa gga gaa aaa cat tat tat tat ata att aag aac	719
Pro Leu Thr Lys Lys Gly Glu Lys His Tyr Tyr Tyr Ile Ile Lys Asn	
225 230 235	

tca tgg gga caa caa tgg gga gaa aga ggt ttc ata aat att gaa aca	767
Ser Trp Gly Gln Gln Trp Gly Glu Arg Gly Phe Ile Asn Ile Glu Thr	
240 245 250 255	

gat gaa tca gga tta atg aga aaa tgt gga tta ggt act gat gca ttc	815
Asp Glu Ser Gly Leu Met Arg Lys Cys Gly Leu Gly Thr Asp Ala Phe	
260 265 270	

att cca tta att gaa cat cat cat cat cat cat taagtcgacg cgatcgaatt	868
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35 40 45

Glu Glu Asn Phe Asp His Ala Ala Tyr Asp Trp Arg Leu His Ser Gly
50 55 60

Val Thr Pro Val Lys Asp Gln Lys Asn Cys Gly Ser Cys Trp Ala Phe
65 70 75 80

Ser Ser Ile Gly Ser Val Glu Ser Gln Tyr Ala Ile Arg Lys Asn Lys  
85 90 95

Leu Ile Thr Leu Ser Glu Gln Glu Leu Val Asp Cys Ser Phe Lys Asn  
100 105 110

Tyr Gly Cys Asn Gly Gly Leu Ile Asn Asn Ala Phe Glu Asp Met Ile  
115 120 125

Glu Leu Gly Gly Ile Cys Pro Asp Gly Asp Tyr Pro Tyr Val Ser Asp  
130 135 140

Ala Pro Asn Leu Cys Asn Ile Asp Arg Cys Thr Glu Lys Tyr Gly Ile  
145 150 155 160

Lys Asn Tyr Leu Ser Val Pro Asp Asn Lys Leu Lys Glu Ala Leu Arg  
165 170 175

Phe Leu Gly Pro Ile Ser Ile Ser Val Ala Val Ser Asp Asp Phe Ala  
180 185 190

Phe Tyr Lys Glu Gly Ile Phe Asp Gly Glu Cys Gly Asp Glu Leu Asn  
195 200 205

His Ala Val Met Leu Val Gly Phe Gly Met Lys Glu Ile Val Asn Pro  
210 215 220

Leu Thr Lys Lys Gly Glu Lys His Tyr Tyr Tyr Ile Ile Lys Asn Ser  
225 230 235 240

Trp Gly Gln Gln Trp Gly Glu Arg Gly Phe Ile Asn Ile Glu Thr Asp  
245 250 255

Glu Ser Gly Leu Met Arg Lys Cys Gly Leu Gly Thr Asp Ala Phe Ile  
260 265 270

Pro Leu Ile Glu His His His His His His  
275 280